

SEQUENCE LISTING

<110> Inbe, Hisayo
Encinas, Jeffrey

<120> Regulation of Human P2Y15 G
Protein-Coupled Receptor

<130> 11582-010-999

<140> 10/529,278

<141> 2005-03-25

<150> PCT/EP2003/010666

<151> 2003-09-25

<150> 60/442,891

<151> 2003-01-28

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<151> 2002-09-27

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Met His Tyr Leu Pro Val Ile Tyr Gly Ile Ile Phe Leu Val Gly Phe	
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Leu Tyr Leu Thr Ser Leu Pro Phe Leu Ile His Tyr Tyr Ala Ser Gly	
85 90 95	

gaa aac tgg atc ttt gga gat ttc atg tgt aag ttt atc cgc ttc agc	336
Glu Asn Trp Ile Phe Gly Asp Phe Met Cys Lys Phe Ile Arg Phe Ser	
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ttc cat ttc aac ctg tat agc agc atc ctc ttc ctc acc tgt ttc agc	384
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Ile Phe Arg Tyr Cys Val Ile Ile His Pro Met Ser Cys Phe Ser Ile	
130 135 140	
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His Lys Thr Arg Cys Ala Val Val Ala Cys Ala Val Val Trp Ile Ile	
145 150 155 160	
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Ser Leu Val Ala Val Ile Pro Met Thr Phe Leu Ile Thr Ser Thr Asn	
165 170 175	
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Arg Thr Asn Arg Ser Ala Cys Leu Asp Leu Thr Ser Ser Asp Glu Leu	
180 185 190	
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Asn Thr Ile Lys Trp Tyr Asn Leu Ile Leu Thr Ala Thr Thr Phe Cys	
195 200 205	
ctc ccc ttg gtg ata gtg aca ctt tgc tat acc acg att atc cac act	672
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210 215 220	
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225 230 235 240	
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Arg Leu Thr Ile Leu Leu Leu Leu Ala Phe Tyr Val Cys Phe Leu Pro	
245 250 255	
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 275 280 285
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Trp Arg Gly Ser Thr Val Ile Met Leu Asn Leu Ala Leu Thr Asp Leu	
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Leu Tyr Leu Thr Ser Leu Pro Phe Leu Ile His Tyr Tyr Ala Ser Gly	
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Glu Asn Trp Ile Phe Gly Asp Phe Met Cys Lys Phe Ile Arg Phe Gly	
100 105 110	
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Phe His Phe Asn Leu Tyr Ser Ser Ile Leu Phe Leu Thr Cys Phe Ser	
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Gln Lys Thr Arg Trp Ala Val Val Ala Cys Ala Gly Val Trp Val Ile	
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Ser Leu Val Ala Val Met Pro Met Thr Phe Leu Ile Thr Ser Thr Thr	
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Arg Thr Asn Arg Ser Ala Cys Leu Asp Leu Thr Ser Ser Asp Asp Leu	
180 185 190	
act act atc aag tgg tac aat ctc att ttg aca gcc acc act ttc tgc	624
Thr Thr Ile Lys Trp Tyr Asn Leu Ile Leu Thr Ala Thr Thr Phe Cys	
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ctg cca ttg gtg ata gtg aca ctt tgc tac acg aca att atc agt acc	672
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225 230 235 240	
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Arg Leu Thr Ile Leu Leu Leu Leu Val Phe Tyr Ile Cys Phe Leu Pro	
245 250 255	
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Phe His Ile Leu Arg Val Ile Arg Ile Glu Ser Arg Leu Leu Ser Ile	
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Pro *	

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 20 25 30
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 Met Gln Tyr Leu Pro Val Ile Tyr Ser Ile Ile Phe Leu Val Gly Phe
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 85 90 95
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 Glu Asn Trp Ile Phe Gly Asp Phe Met Cys Lys Phe Ile Arg Phe Gly
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 Ser Leu Val Ala Val Met Pro Met Thr Phe Leu Ile Thr Ser Thr Thr
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Thr	Thr	Ile	Lys	Trp	Tyr	Asn	Leu	Ile	Leu	Thr	Ala	Thr	Thr	Phe	Cys	
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Leu	Pro	Leu	Leu	Ile	Val	Thr	Leu	Cys	Tyr	Thr	Thr	Ile	Ile	Ser	Thr	
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Pro	Leu	Ala	Ala	Leu	Asn	Thr	Phe	Gly	Asn	Leu	Leu	Leu	Tyr	Val	Val	
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Val	Ser	Asn	Asn	Phe	Gln	Gln	Ala	Phe	Cys	Ser	Ala	Val	Arg	Cys	Lys	
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Ala	Ile	Gly	Asp	Leu	Glu	Gln	Ala	Lys	Lys	Asp	Ser	Cys	Ser	Asn	Asn	
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Pro	*															

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